AMENDMENTS TO SPECIFICATION

The additions have been indicated by underlining.

Please replace the paragraph which begins as the fourth paragraph on page 29 and continues on to the top of page 30 with the following amended paragraph:

To evaluate randomisation of the region between the start codon and the gene of interest, a vector library can be designed consisting of a randomised stretch of five amino acids adjacent to the start codon. This creates a great vector variation and ideally the randomised region can be fitted individually to the different protein targets. The randomised nucleotides after the ATG in the expression vector can either be added to the vector directly or added together with the gene of interest by cloning of a PCR product generated from randomised primers. The principle of primer design for adding randomised nucleotides is ATG NNN NNN NNN NNN NNN [SEQ ID NO: 1]. The selection of highly expressing soluble clones will be performed as described with the CoFi-blot technique.

Attachment: Sequence Listing